

FIG. 2

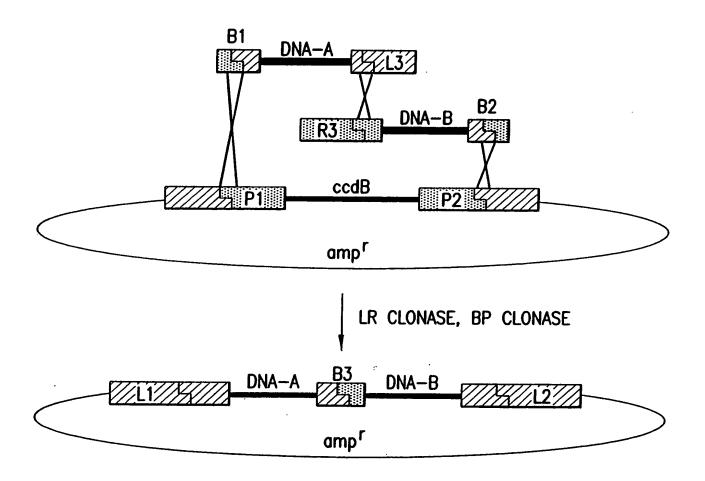


FIG. 3

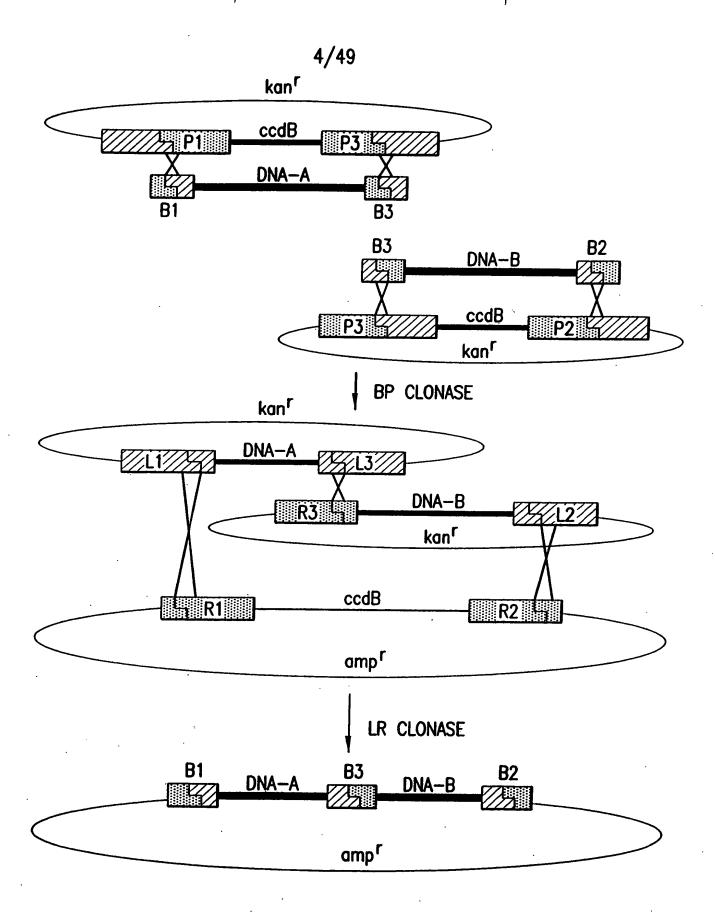


FIG. 4

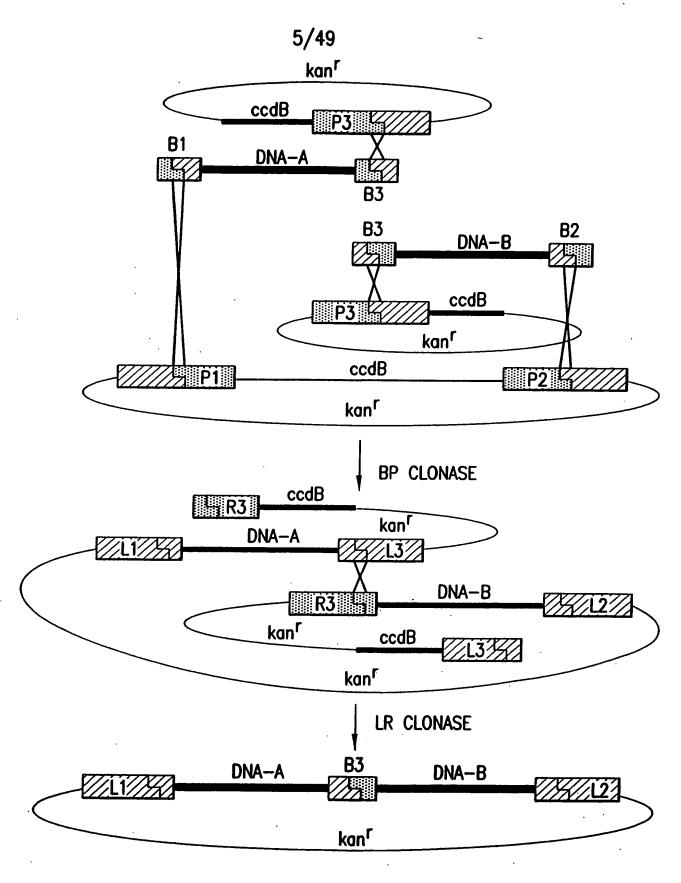


FIG. 5

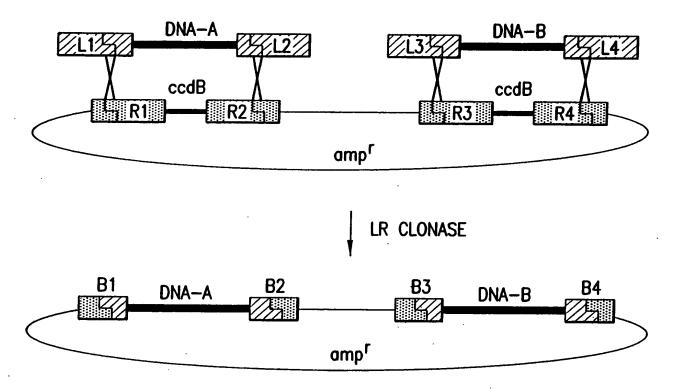


FIG. 6

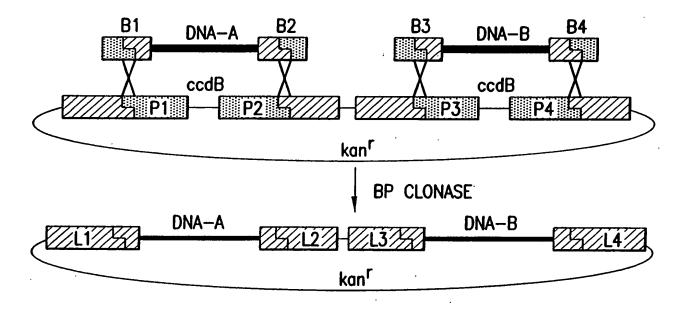
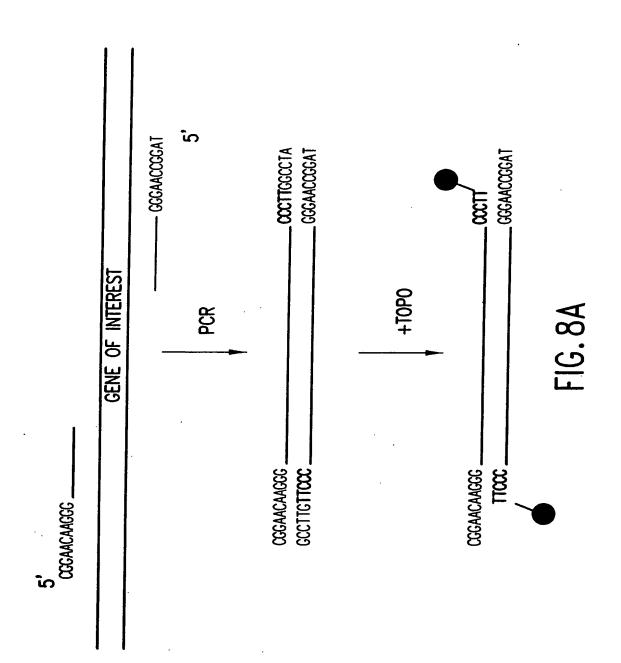


FIG. 7



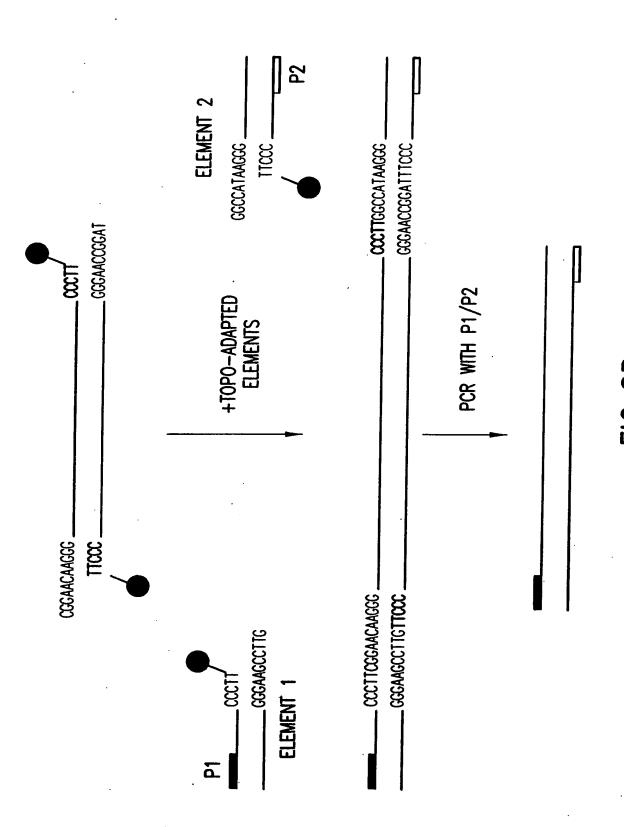


FIG. 8B

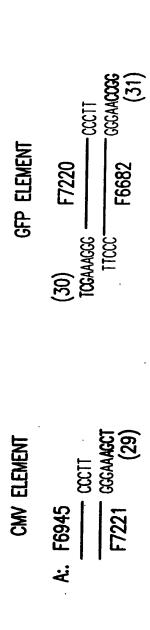
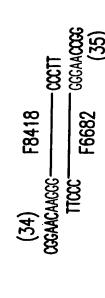


FIG. 9A



F8417

B. F6945



F8417

FIG. 9C

BGH ELEMENT	F7222	F6948
	32) 360CAAGGG	

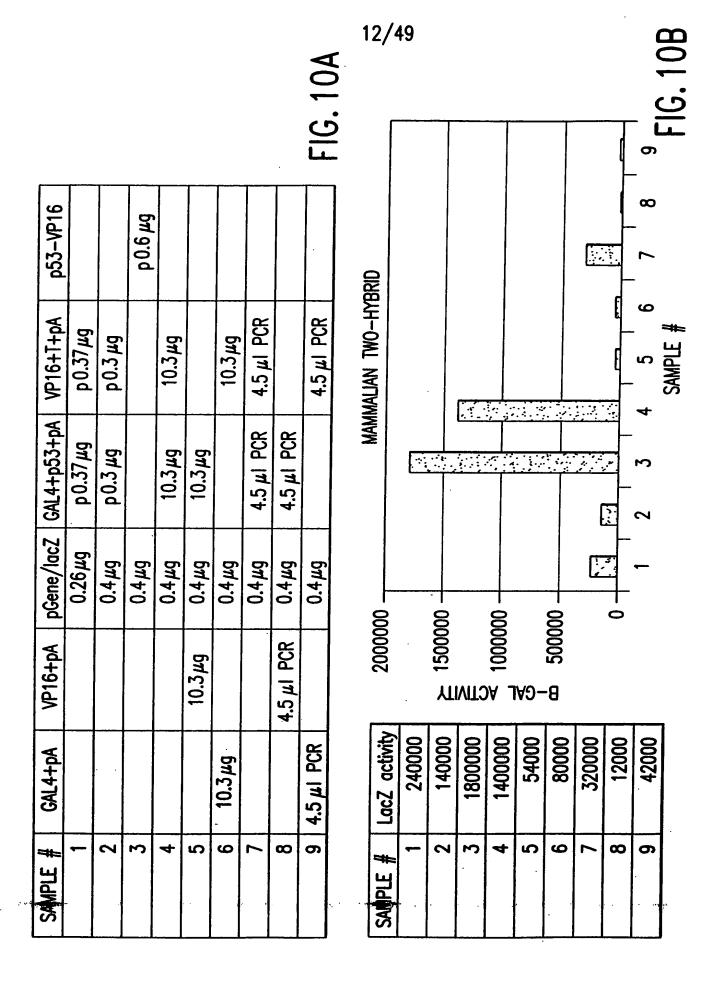
F7222		F6948
(36)	GGCCAAGG(

F8419	8	TTCC FROAR
(40)	GCCTAAAGGC-	211

TABLE 1

Primer name	F#	Sequence (5'-> 3')	SEQ ID NO:
MTH1	10779	TATGTATCATACACATACGATTTAGGT	1
MTH2	10780	ACOGCCTCTCCCCGCGCGTT	2
GAL4r2	12667	GTTCCGAAGGGGGGATACAGTCAACTGTCTTTG	3
MTH5	12505	TTGGCCAAGGGTATCTAGAAGCTTCTGCAGACGCGT	4
VP16r2	12668	GTTCCGAAGGGCCACCGTACTCGTCAATTCCAAG	1 5
SV40pAf	12016	GGCCAAAAGGGAACTTGTTTATTGCAGCTTATAATG	6
SV40pAr	561	CTCTGACTTGAGCGTCGATTTT	7
p53f2	12669	CGGAACAAGGGGAATTCCCTGTCACCGAGACC	8
SVTf2	12670	CGGAACAAGGGGAATTCCCGGGGATCTGGAATTC	9
CMVr2	7221	TCGAAAGGCTCGAGCTGCAGCTG	10
CMVf	6945	AATTCACATTGATTATTGAGTAGTTA	11
GFP-Xhof	7220	TOGAAAGGTAATGGCCAGCAAAGGAGAAG	12
GFP-Notr	6682	GGCCAAGGGTTTGTAGAGCTCATCCAT	13
BGHf2	7222	GGCCAAGGGTCTGAATGGGGCCGCATAGT	14
BGHr	6948	AAGCCATAGAGCCCGGGCCA	15
CMVr3	8417	GTTCCGAAGGGTCGAGCTGCAGCTG	16
GFPf3	8418	CGGAACAAGGATGGCCAGCAAAGGAGAAG	17
GFPr3	8420	TAGGCCAAGGGTTTGTAGAGCTCATCCATGC	18
BGHf3	8419	GGCCTAAAGGGTGAATGGGGCCGCATAGT	19
T7top	9304	GAAGGAGTAATACGACTCACTATAGGGAGCCACCATGGGCCCTTCGGAAC	20
T7bottom	9305	GTTCCGAAGGCCCCATGGTGGCTCCCTATAGTGAGTCGTATTACTCCTTC	21
T7amp	9306	GAAGGAGTAATACGACTCACT	22
T3top	9661	GGCCTAAAGGGTCCCTTTAGTGAGGGTTAATTGCGCGC	23
T3bottom	9662	GCGCGCAATTAACCCTCACTAAAGGGACCCTTTAGGCC	24
lacZf2	10632	CGGAACAAGGGATGATAGATCCCGTCGTTTTACA	25
lacZ1k2	10770	TACGCCAAGGGGACCATTTTCAATCCGCACCT	26
lacZ2k2	10771	TAGGCCAAGGGGAGGCACTTCACCGCTTGCCA	27
lacZ3k2	10772	TAGGCCAAGGGTTTGACACCAGACCAACTGGTA	28

FIG. 9D



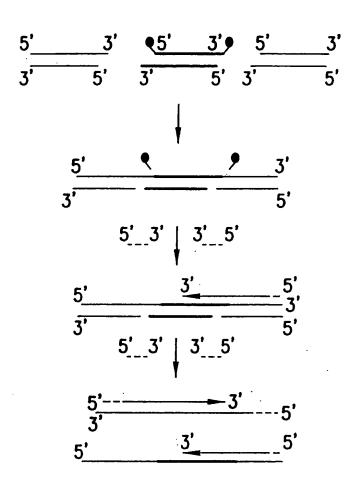
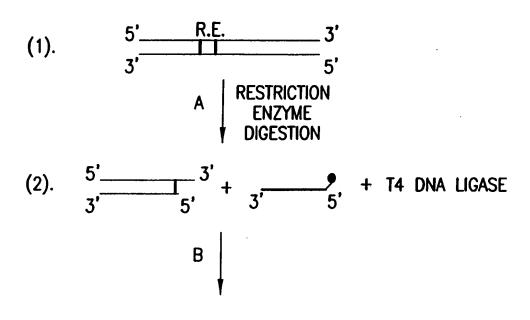


FIG. 13



(3).
$$\frac{5'}{3'}$$
 + $\frac{3'}{5'}$ + $\frac{3'}{5'}$ + T4 DNA LIGASE

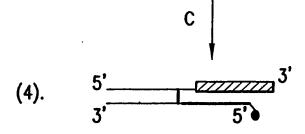


FIG. 14

SS DNA
$$\frac{3}{5}$$
, + SSRNA $\frac{9}{3}$, 5', $\frac{9}{5}$, $\frac{3}{5}$, $\frac{9}{5}$, $\frac{3}{5}$, $\frac{3}$, $\frac{3}{5}$, $\frac{3}{5}$, $\frac{3}{5}$, $\frac{3}{5}$, $\frac{3}{5}$, $\frac{3}$

FIG. 15

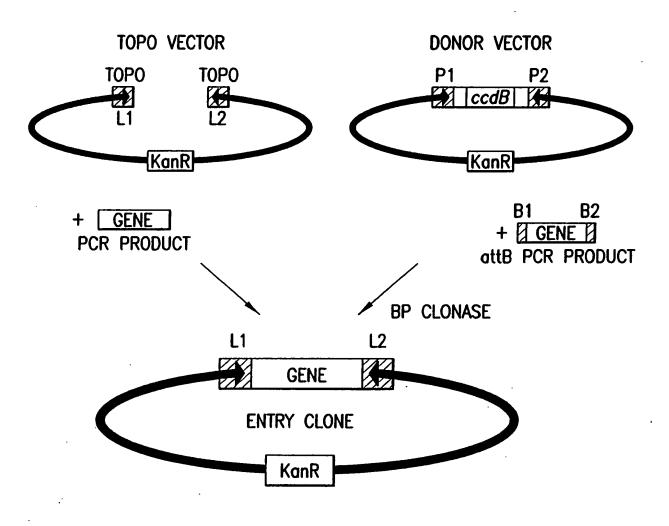
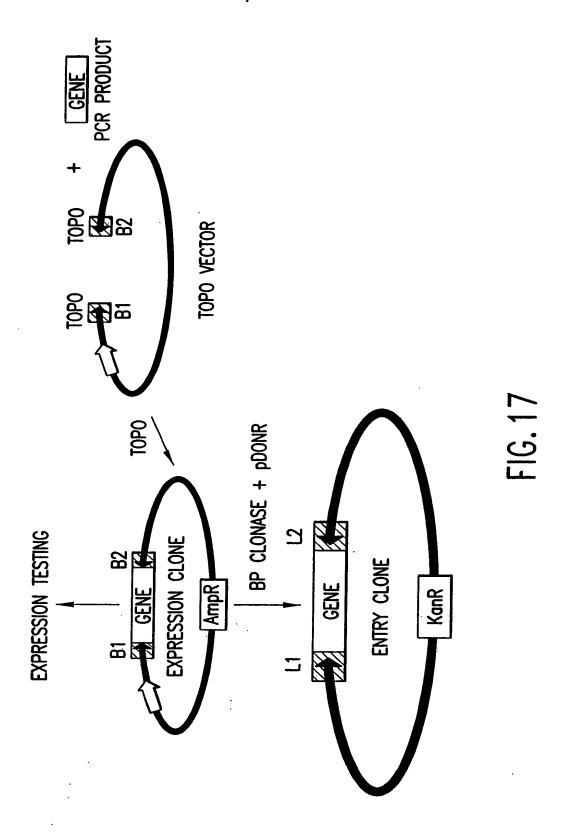


FIG. 16



MCS FOR pcDNAGW-DT(sc) AND pENTR-DT(sc)

>	Ę	5		
×	AAA	إ		
>	TAC	: ا	7	
_	13		3	
4	TC	:		82
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«	IIG IAC AMA AMA GCA GGC TOC GCG GCC GCA CTC GAG AMA GGC GGC GGC GCC CCA GCT TTC TTC TAC AMA CTC			
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FIG. 18

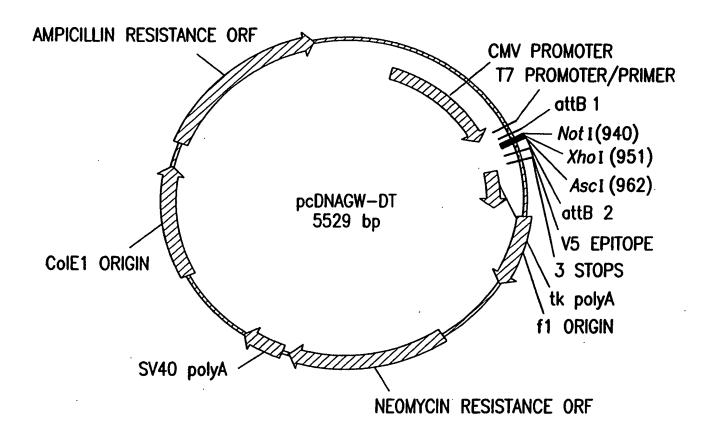
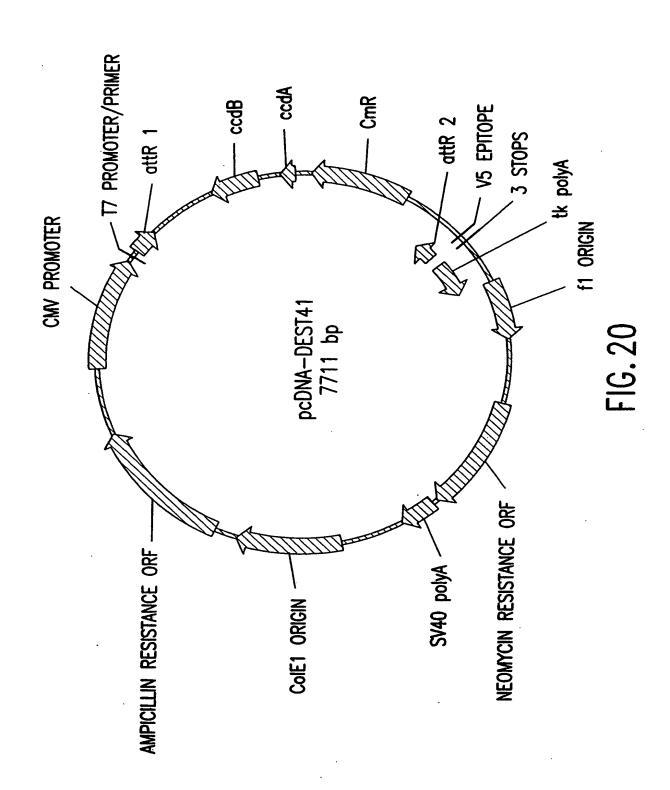
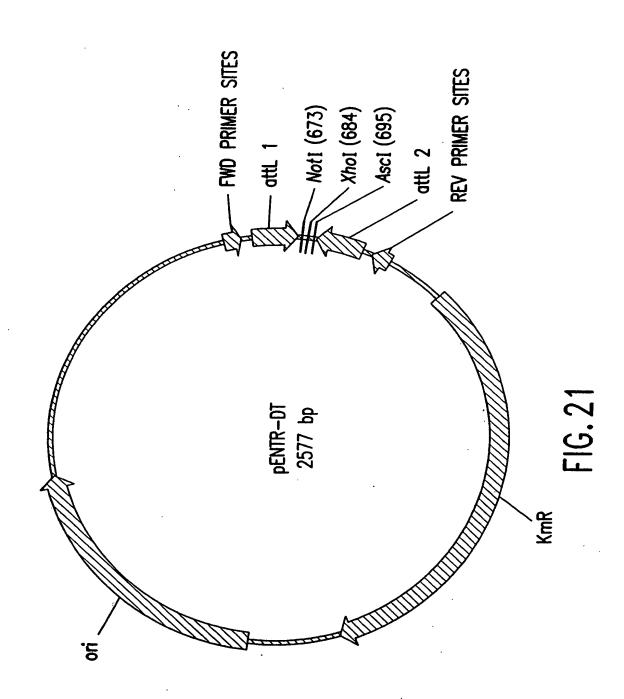


FIG. 19





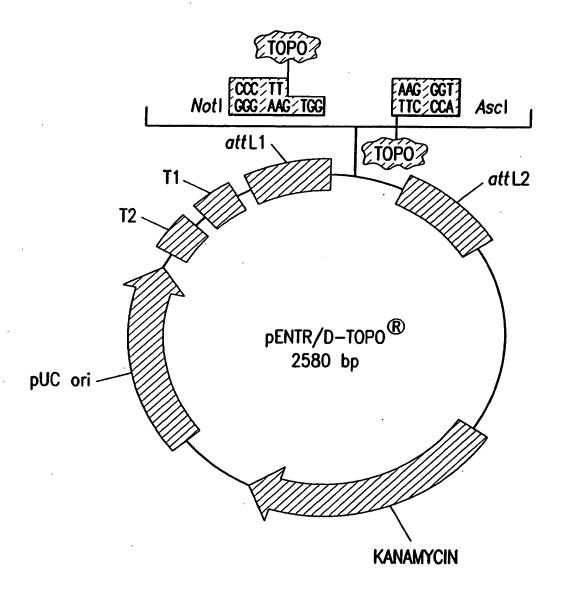


FIG. 22A

1 ctttcctgcg ttatcccctg attctgtgga taaccgtatt accgcctttg agtgagctga 61 taccgctcgc cgcagccgaa cgaccgagcg cagcgagtca gtgagcgagg aagcggaaga 121 gcgcccaata cgcaaaccgc ctctccccgc gcgttggccg attcattaat gcagctggca 181 cgacaggttt cccgactgga aagcgggcag tgagcgcaac gcaattaata cgcgtaccgc 241 tagccaggaa gagtttgtag aaacgcaaaa aggccatccg tcaggatggc cttctgctta 301 gtttgatgcc tggcagttta tggcgggcgt cctgcccgcc accctccggg ccgttgcttc 361 acaacgttca aatccgctcc cggcggattt gtcctactca ggagagcgtt caccgacaaa 421 caacagataa aacgaaaggc ccagtcttcc gactgagcct ttcgttttat ttgatgcctg 481 gcagttccct actctcgcgt taacgctagc atggatgttt tcccagtcac gacgttgtaa 541 aacgacggcc agtcttaagc tcgggcccca aataatgatt ttattttgac tgatagtgac 601 ctgttcgttg caacaaattg atgagcaatg cttttttata atgccaactt tgtacaaaaa 661 agcaggetee geggeegee etteaceatg nnnnnnnna agggtgggeg egeegaeeea 721 gctttcttgt acaaagttgg cattataaga aagcattgct tatcaatttg ttgcaacgaa 781 caggicacta tcagicaaaa taaaatcatt attigccatc cagcigatat cccctatagi 841 gagtcgtatt acatggtcat agctgtttcc tggcagctct ggcccgtgtc tcaaaatctc 901 tgatgttaca ttgcacaaga taaaaatata tcatcatgaa caataaaact gtctgcttac 961 ataaacagta atacaagggg tgttatgagc catattcaac gggaaacgtc gaggccgcga 1021 ttaaattcca acatggatgc tgatttatat gggtataaat gggctcgcga taatgtcggg 1081 caatcaggtg cgacaatcta tcgcttgtat gggaagcccg atgcgccaga gttgtttctg 1141 aaacatggca aaggtagcgt tgccaatgat gttacagatg agatggtcag actaaactgg 1201 ctgacggaat ttatgcctct tccgaccatc aagcatttta tccgtactcc tgatgatgca 1261 tggttactca ccactgcgat ccccggaaaa acagcattcc aggtattaga agaatatcct 1321 gattcaggtg aaaatattgt tgatgcgctg gcagtgttcc tgcgccggtt gcattcgatt 1381 cctgtttgta attgtccttt taacagcgat cgcgtatttc gtctcgctca ggcgcaatca 1441 cgaatgaata acggtttggt tgatgcgagt gattttgatg acgagcgtaa tggctggcct 1501 gttgaacaag totggaaaga aatgcataaa ottttgccat totcacogga ttcagtogto 1561 actcatggtg atttctcact tgataacctt atttttgacg aggggaaatt aataggttgt 1621 attgatgttg gacgagtcgg aatcgcagac cgataccagg atcttgccat cctatggaac 1681 tgcctcggtg agttttctcc ttcattacag aaacggcttt ttcaaaaata tggtattgat 1741 aatcctgata tgaataaatt gcagtttcat ttgatgctcg atgagttttt ctaatcagaa 1801 ttggttaatt ggttgtaaca ctggcagagc attacgctga cttgacggga cggcgcaagc 1861 tcatgaccaa aatcccttaa cgtgagttac gcgtcgttcc actgagcgtc agaccccgta 1921 gaaaagatca aaggatcttc ttgagatcct ttttttctgc gcgtaatctg ctgcttgcaa 1981 acaaaaaac caccgctacc agcggtggtt tgtttgccgg atcaagagct accaactctt 2041 tttccgaagg taactggctt cagcagagcg cagataccaa atactgtcct tctagtgtag 2101 ccgtagttag gccaccactt caagaactct gtagcaccgc ctacatacct cgctctgcta 2161 atcctgttac cagtggctgc tgccagtggc gataagtcgt gtcttaccgg gttggactca 2221 agacgatagt taccggataa ggcgcagcgg tcgggctgaa cggggggttc gtgcacacag 2281 cccagcttgg agcgaacgac ctacaccgaa ctgagatacc tacagcgtga gcattgagaa 2341 agcgccacgc ttcccgaagg gagaaaggcg gacaggtatc cggtaagcgg cagggtcgga 2401 acaggagage geacgaggga gettecaggg ggaaaegeet ggtatettta tagteetgte 2461 gggtttcgcc acctctgact tgagcgtcga tttttgtgat gctcgtcagg ggggcggagc 2521 ctatggaaaa acgccagcaa cgcggccttt ttacggttcc tggccttttg ctggcctttt 2581 gctcacatgt t

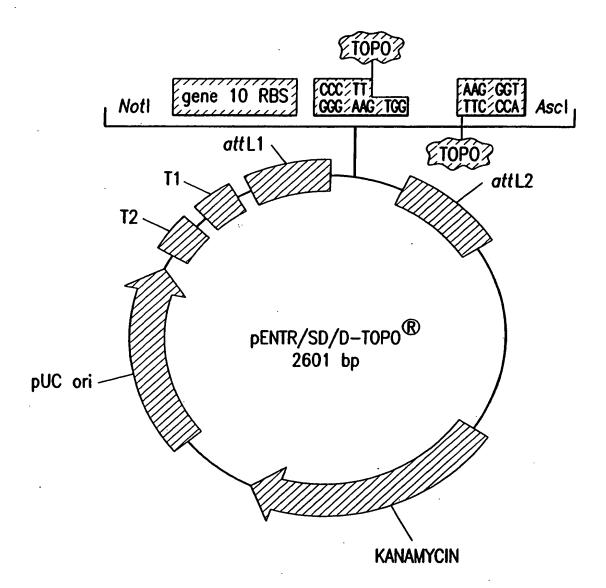
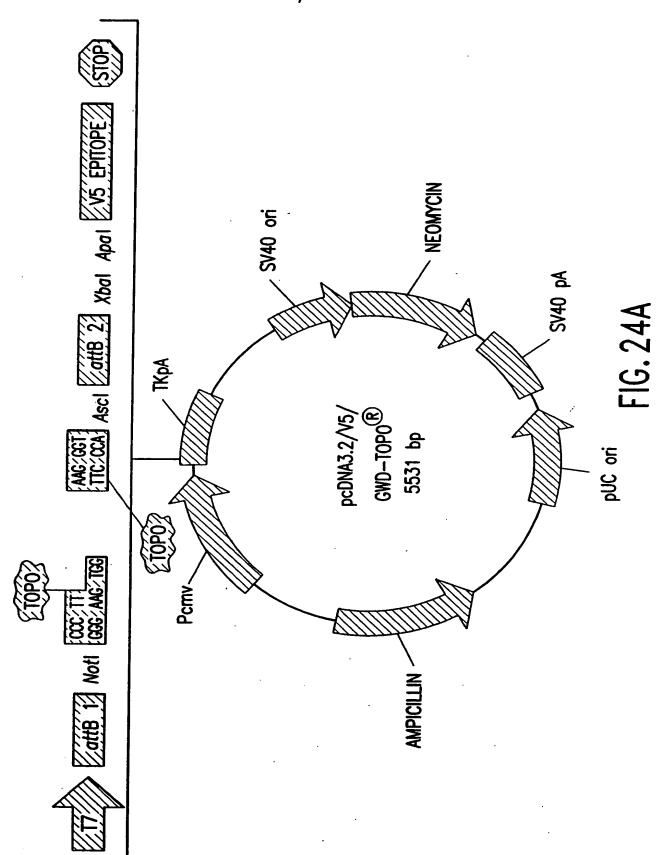


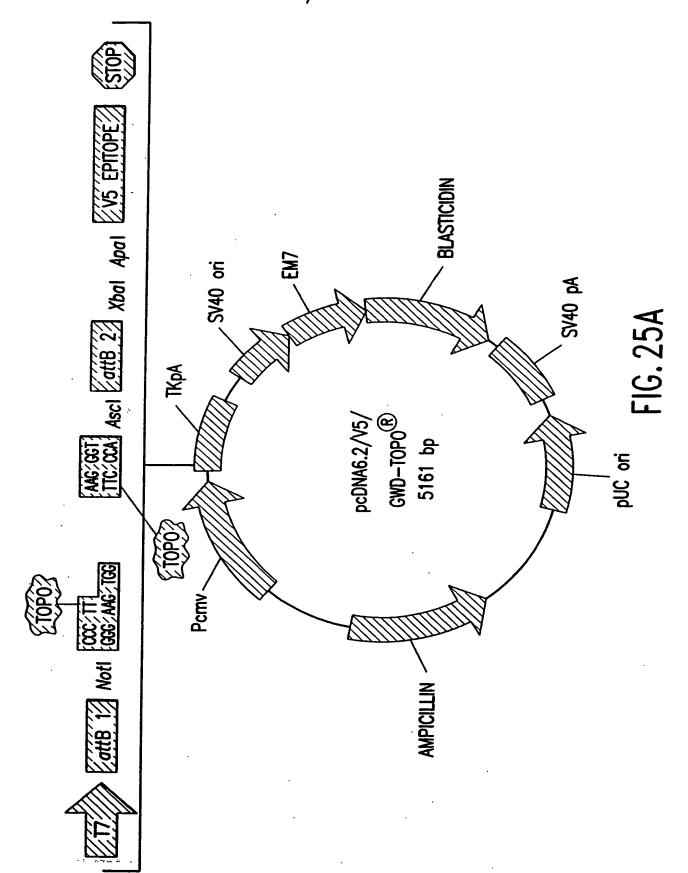
FIG. 23A

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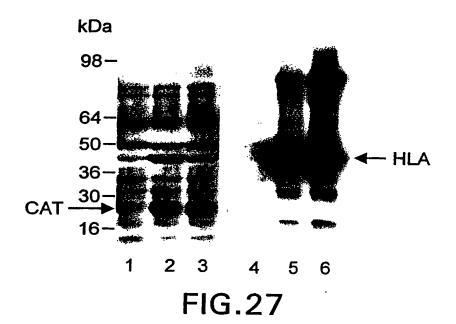


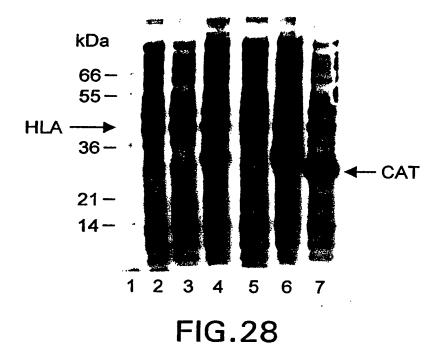
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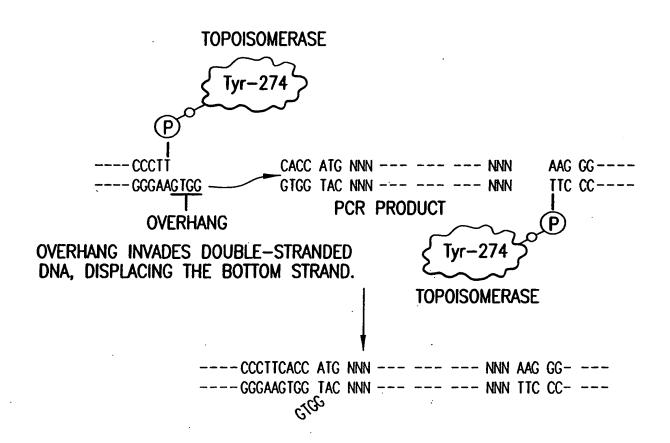
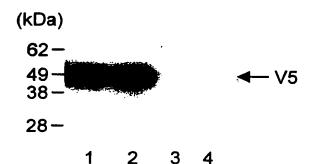
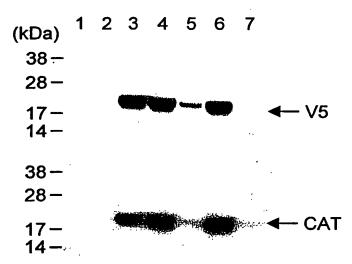


FIG. 29



Lane 1: pCMVTetO/CAT/V5TKpA (without secondary PCR) + Tet Lane 2: pCMVTetO/CAT/V5TKpA (with secondary PCR) + Tet Lane 3: pCMVTetO/CAT/V5TKpA (with secondary PCR) - Tet Lane 4: pCMVTetO/CAT/V5TKpA (without secondary PCR) - Tet

FIG.30A



Lane 1: TRex-CHO Cells + Tet

Lane 2: without secondary PCR (with purified CAT) - Tet

Lane 3: without secondary PCR (with purified CAT) + Tet

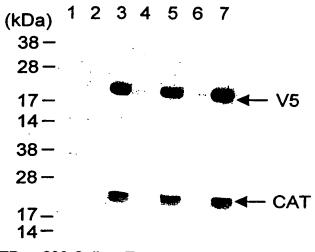
Lane 4: without secondary PCR (with unpurified CAT) + Tet

Lane 5: without secondary PCR (with unpurified CAT) -Tet

Lane 6: with secondary PCR + Tet

Lane 7: with secondary PCR - Tet

FIG.30B



Lane 1: TRex-293 Cells + Tet

Lane 2: without secondary PCR (with purified CAT) - Tet

Lane 3: without secondary PCR (with purified CAT) + Tet

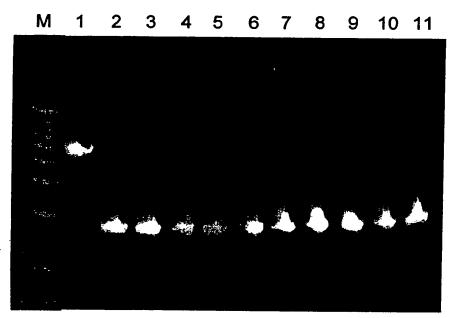
Lane 4: without secondary PCR (with unpurified CAT) - Tet

Lane 5: without secondary PCR (with unpurified CAT) +Tet

Lane 6: with secondary PCR - Tet

Lane 7: with secondary PCR + Tet

FIG.30C



Lane 1: negative control; lanes 2-11: test clones; M: 500 bp marker

FIG.31

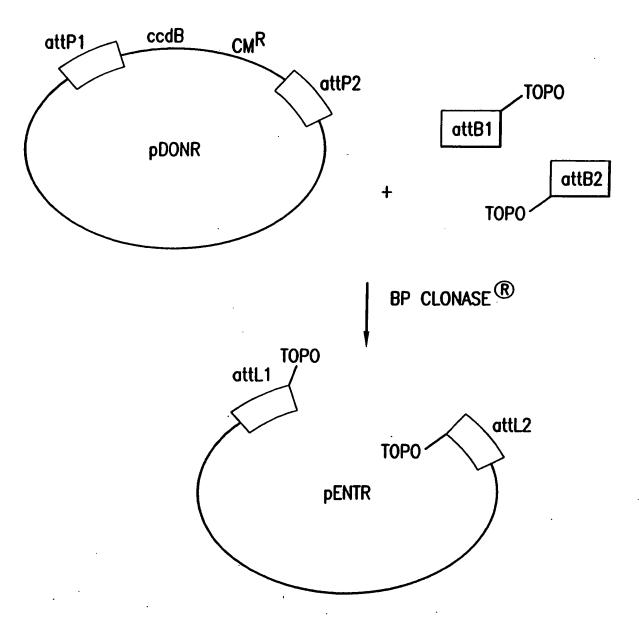
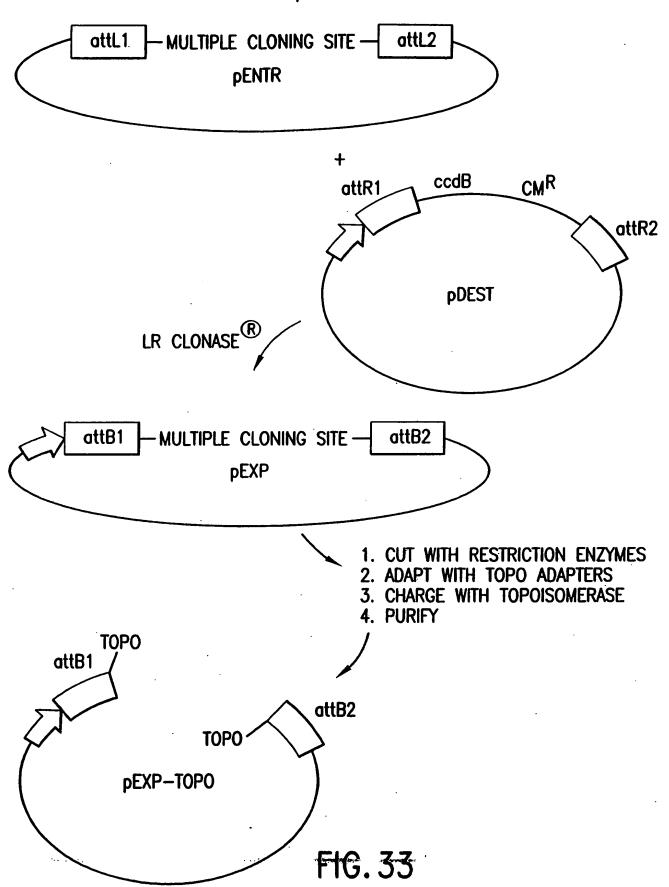


FIG. 32



ASSEMBLE NUCLEIC ACID SEGMENTS

AMPLIFY ASSEMBLED
NUCLEIC ACID
SEGMENTS

USE ASSEMBLED
NUCLEIC ACID
SEGMENTS

USE AMPLIFIED
ASSEMBLED NUCLEIC
ACID SEGMENTS

FIG. 34

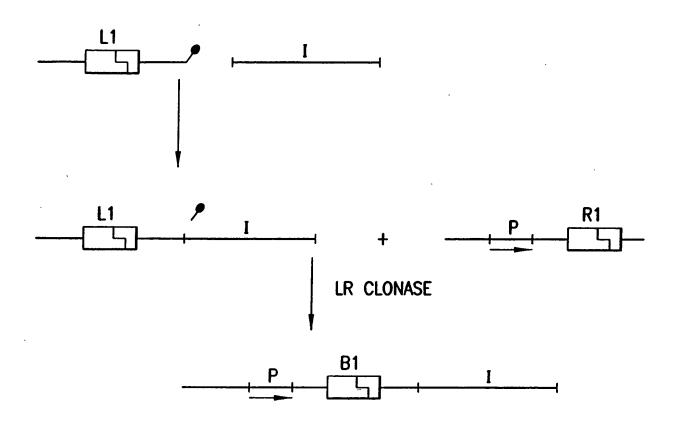
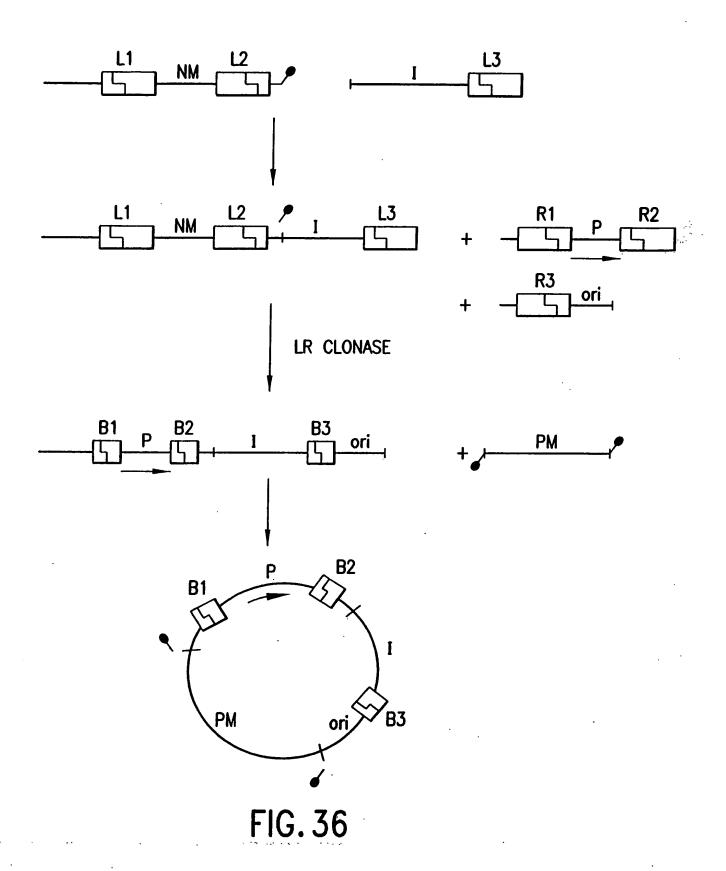


FIG. 35



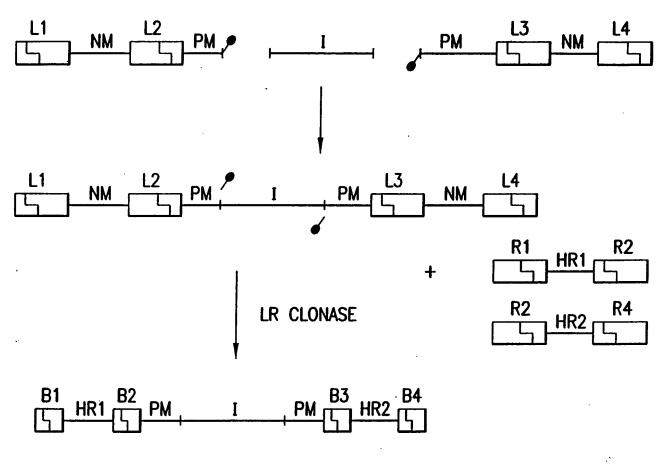


FIG. 37

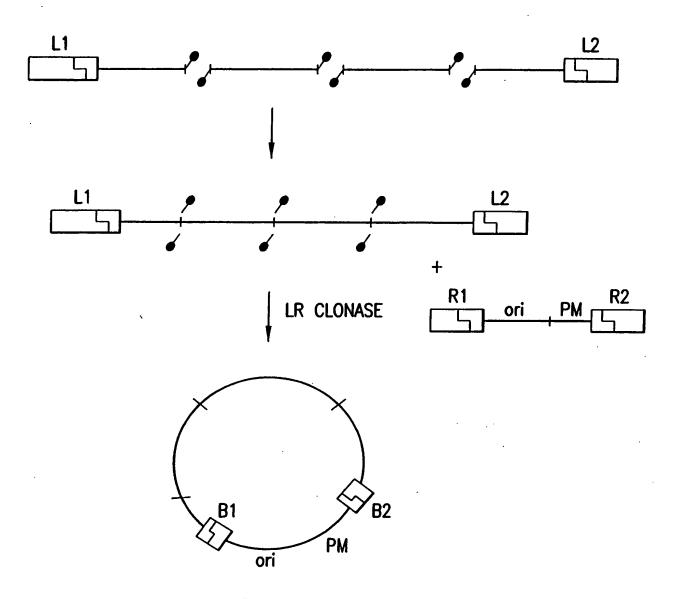


FIG. 38

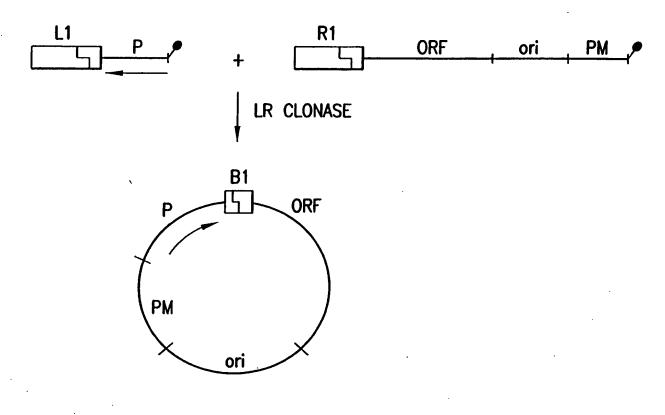


FIG. 39

FIG. 40

Figure 41 PER PRODUCT S' T7 PROMOTER TH LIGHSE PROMOTER PCR PRODUCT T 7
PROMOTER AMPLIFICATION USINA PRIMERS A AND B PCR PRODUCT

T7 promoter TOPO

5' pGACTCGTAATACGACTCACTATAGGGCCCTT 3'

3' AAAAAAAAAAAACTGAGCATTATGCTGAGTGATATCCCGGGAp 5'

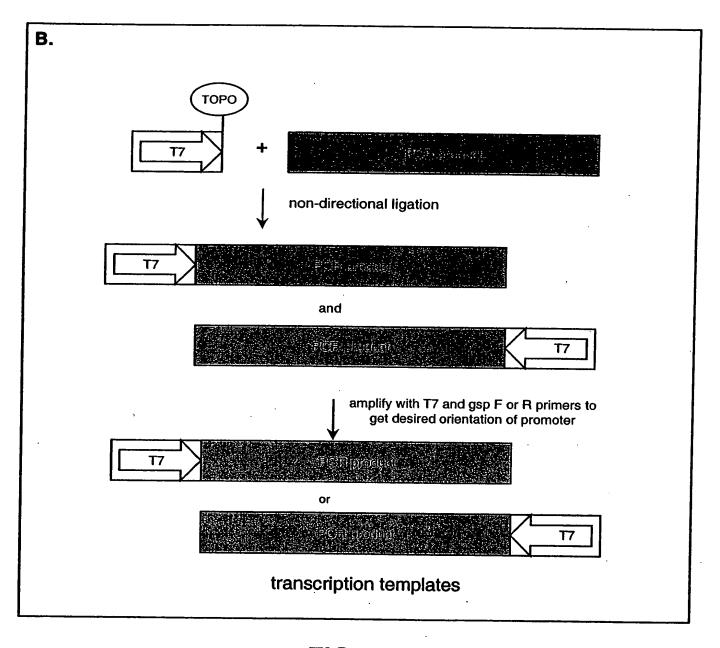
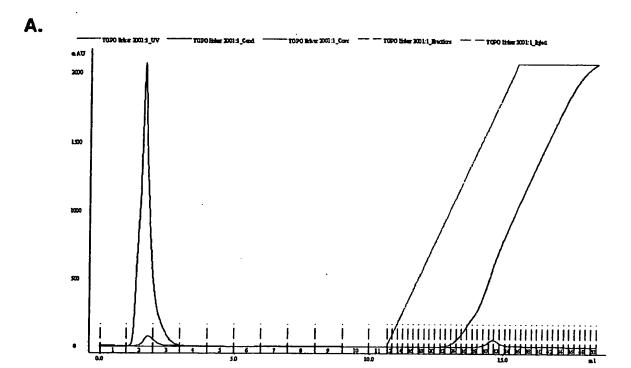


FIG. 42



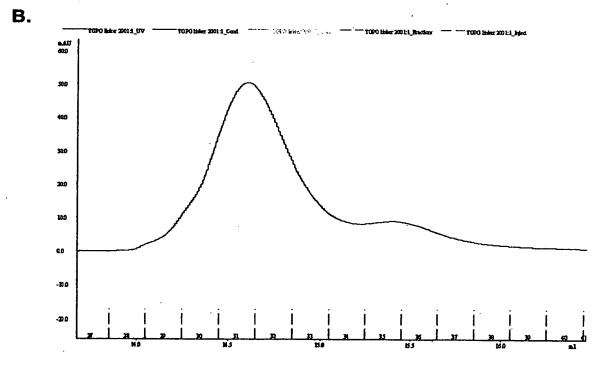
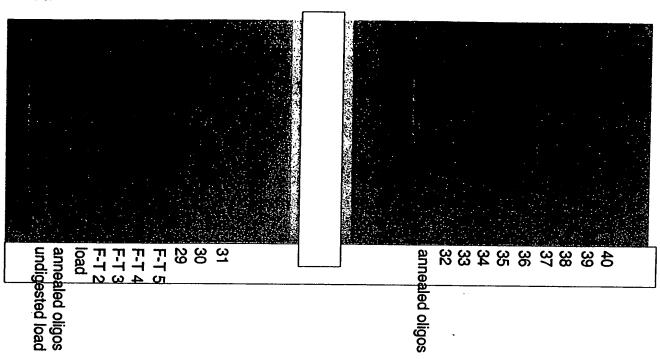


FIG. 43





B.

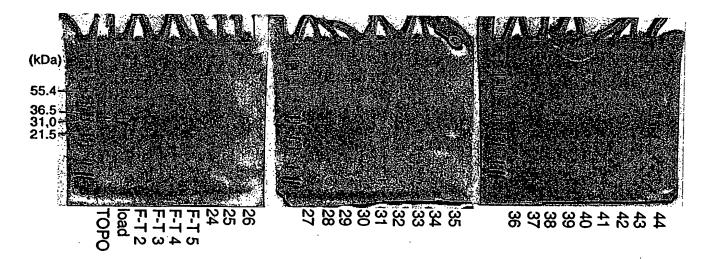


FIG. 44

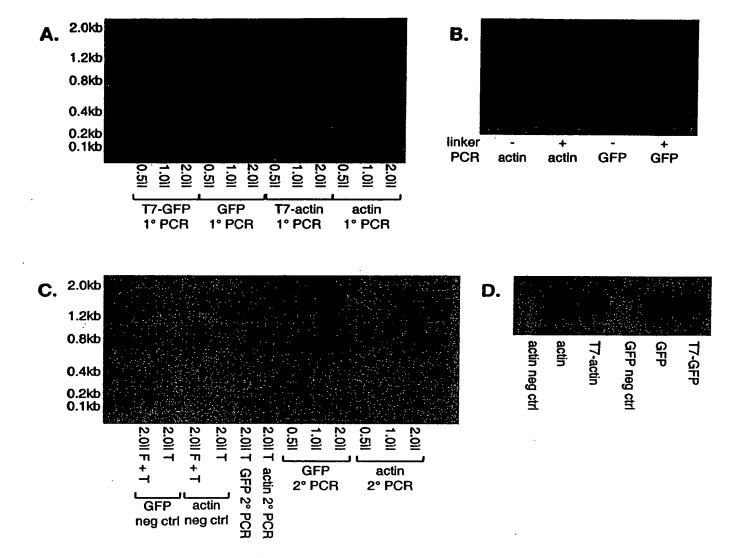


FIG. 45

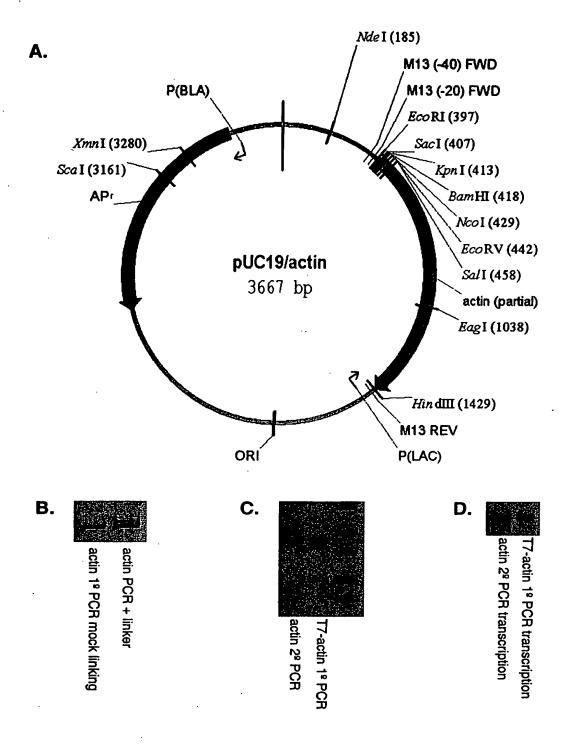


FIG. 46